RAW SEQUENCE LISTING

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Application Serial Number: 10/049 404
Source: PCT
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RAW SEQUENCE LISTING DATE: 05/24/2005
PATENT APPLICATION: US/10/049,404 TIME: 15:03:57

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<110> APPLICANT: Arndt, Michaela
Little, Melvyn
Kypriyanov, Sergey
Krauss, Jurgen
Pfreundschuh, Michael

<120> TITLE OF INVENTION: Fv Antibody Construct Comprising Binding Sites For a CD16 Receptor and a
CD30 Surface Protein

<130> FILE REFERENCE: 4121-135

<140> CURRENT APPLICATION NUMBER: US 10/049,404

<141> CURRENT FILING DATE: 2002-02-05

<150> PRIOR APPLICATION NUMBER: PCT/DE00/02589

<151> PRIOR FILING DATE: 2000-08-02

<150> PRIOR APPLICATION NUMBER: DE 199 37 264

<151> PRIOR FILING DATE: 1999-08-06

<160> NUMBER OF SEQ ID NOS: 11

<170> SOFTWARE: PatentIn version 3.1

<210> SEQ ID NO: 1

<211> LENGTH: 4570

<212> TYPE: DNA

<213> ORGANISM: Artificial Sequence

<220> FEATURE:

<223> OTHER INFORMATION: Synthetic Construct

<400> SEQUENCE: 1

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206             20                  25                  30
209 Asn Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His Gly Leu Glu
210         35                  40                  45
213 Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn Tyr Asn Glu
214     50                  55                  60
217 Lys Phe Lys Gly Lys Ala Thr Val Thr Ala Asp Thr Ser Ser Arg Thr
218 65                  70                  75                  80
221 Ala Tyr Val Gln Val Arg Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
222                 85                  90                  95
225 Phe Cys Ala Arg Ser Ala Ser Trp Tyr Phe Asp Val Trp Gly Ala Arg
226             100                 105                 110
229 Thr Thr Val Thr Val Ser Ser Ala Lys Thr Thr Pro Lys Leu Gly Gly
230         115                 120                 125
233 Asp Ile Glu Leu Thr Gln Ser Pro Lys Phe Met Ser Thr Ser Val Gly
234     130                 135                 140
237 Asp Arg Val Asn Val Thr Tyr Lys Ala Ser Gln Asn Val Gly Thr Asn
238 145                 150                 155                 160
241 Val Ala Trp Phe Gln Gln Lys Pro Gly Gln Ser Pro Lys Val Leu Ile
242                 165                 170                 175
245 Tyr Ser Ala Ser Tyr Arg Tyr Ser Gly Val Pro Asp Arg Phe Thr Gly
246             180                 185                 190
249 Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Asn Val Gln Ser
250         195                 200                 205
253 Glu Asp Leu Ala Glu Tyr Phe Cys Gln Gln Tyr His Thr Tyr Pro Leu
254     210                 215                 220
257 Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala
258 225                 230                 235                 240
261 Ala Ala Gly Ser Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser
262                 245                 250                 255
265 His His His His His His
266             260
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287 Thr Tyr Thr Ile His Trp Val Arg Gln Arg Pro Gly His Asp Leu Glu
288         35                  40                  45
291 Trp Ile Gly Tyr Ile Asn Pro Ser Ser Gly Tyr Ser Asp Tyr Asn Gln
292     50                  55                  60
295 Asn Phe Lys Gly Lys Thr Thr Leu Thr Ala Asp Lys Ser Ser Asn Thr
296 65                  70                  75                  80
299 Ala Tyr Met Gln Leu Asn Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr
300                 85                  90                  95
303 Tyr Cys Ala Arg Arg Ala Asp Tyr Gly Asn Tyr Glu Tyr Thr Trp Phe
304             100                 105                 110
307 Ala Tyr Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Lys Thr
308         115                 120                 125
311 Thr Pro Lys Leu Gly Gly Asp Ile Gln Ala Val Val Thr Gln Glu Ser
312     130                 135                 140
315 Ala Leu Thr Thr Ser Pro Gly Glu Thr Val Thr Leu Thr Cys Arg Ser
316 145                 150                 155                 160
319 Asn Thr Gly Thr Val Thr Thr Ser Asn Tyr Ala Asn Trp Val Gln Glu
320                 165                 170                 175
323 Lys Pro Asp His Leu Phe Thr Gly Leu Ile Gly His Thr Asn Asn Arg
324             180                 185                 190
327 Ala Pro Gly Val Pro Ala Arg Phe Ser Gly Ser Leu Ile Gly Asp Lys
328         195                 200                 205
331 Ala Ala Leu Thr Ile Thr Gly Ala Gln Thr Glu Asp Glu Ala Ile Tyr
332     210                 215                 220
335 Phe Cys Ala Leu Trp Tyr Asn Asn His Trp Val Phe Gly Gly Gly Thr
336 225                 230                 235                 240
339 Lys Leu Thr Val Leu Gly Gln Pro Lys Ser Ala Ala Ala Gly Ser Glu
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400 <211> LENGTH: 21
401 <212> TYPE: DNA
402 <213> ORGANISM: Artificial Sequence
404 <220> FEATURE:
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407 <400> SEQUENCE: 8
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412 <211> LENGTH: 36
413 <212> TYPE: DNA
414 <213> ORGANISM: Artificial Sequence
416 <220> FEATURE:
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424 <211> LENGTH: 27
425 <212> TYPE: DNA
426 <213> ORGANISM: Artificial Sequence
428 <220> FEATURE:
429 <223> OTHER INFORMATION: Synthetic Construct
431 <400> SEQUENCE: 10
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 05/24/2005
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Input Set : A:\EP.txt
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The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 9

VERIFICATION SUMMARY DATE: 05/24/2005
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Input Set : A:\RP.txt
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